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TECH CENTER 1600/2900

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RAW SEQUENCE LISTING
PATENT APPLICATION: **US/08/599,974E**

DATE: 02/11/2003
TIME: 10:56:27

Input Set : **A:\EP.txt**
Output Set: **N:\CRF4\02112003\H599974E.raw**

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 3 <110> APPLICANT: The Rockefeller University
 4        Friedman, Jeffrey M.
 5        Lee, Gwo-Hwa
 6        Proenca, Ricardo
 8 <120> TITLE OF INVENTION: DB, THE RECEPTOR FOR LEPTIN,NUCLEIC ACIDS ENCODING THE
RECEPTOR, AND USES
 9        THEREOF
11 <130> FILE REFERENCE: 600-1-162CP1
13 <140> CURRENT APPLICATION NUMBER: 08/599,974E
14 <141> CURRENT FILING DATE: 1996-02-14
16 <150> PRIOR APPLICATION NUMBER: US 09/586,594
17 <151> PRIOR FILING DATE: 1996-01-16
19 <160> NUMBER OF SEQ ID NOS: 97
21 <170> SOFTWARE: PatentIn version 3.1
23 <210> SEQ ID NO: 1
24 <211> LENGTH: 2529
25 <212> TYPE: DNA
26 <213> ORGANISM: Mus musculus
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33 aatgagcaag gtcaaaactg ctctgcactc acagacaaca ctgaagggaa gacactggct     180
35 tcagtagtga aggcttcagt ttttcgccag ctaggtgtaa actgggacat agagtgctgg     240
37 atgaaagggg acttgacatt attcatctgt catatggagc cattacctaa gaaccccttc     300
39 aagaattatg actctaaggt ccatctttta tatgatctgc ctgaagtcat agatgattcg     360
41 cctctgcccc cactgaaaga cagctttcag actgtccaat gcaactgcag tcttcgggga     420
43 tgtgaatgtc atgtgccggt acccagagcc aaactcaact acgctcttct gatgtatttg     480
45 gaaatcacat ctgccggtgt gagttttcag tcacctctga tgtcactgca gcccatgctt     540
47 gttgtgaaac ccgatccacc cttaggtttg catatggaag tcacagatga tggtaattta     600
49 aagatttctt gggacagcca acaatggca ccatttccgc ttcaatatca ggtgaaatat     660
51 ttagagaatt ctacaattgt aagagaggct gctgaaattg tctcagctac atctctgctg     720
53 gtagacagtg tgcttcctgg atcttcatat gaggtccagg tgaggagcaa gagactggat     780
55 ggttcaggag tctggagtga ctggagttca cctcaagtct ttaccacaca agatgttgtg     840
57 tattttccac ccaaaattct gactagtgtt ggatcgaatg cttcttttca ttgcatctac     900
59 aaaaacgaaa accagattat ctcctcaaaa cagatagttt ggtggaggaa tctagctgag     960
61 aaaatccctg agatacagta cagcattgtg agtgaccgag ttagcaaagt taccttctcc    1020
63 aacctgaaag ccaccagacc tcgagggaag tttacctatg acgcagtgta ctgctgcaat    1080
65 gagcaggcgt gccatcaccg ctatgctgaa ttatacgtga tcgatgtcaa tatcaatata    1140
67 tcatgtgaaa ctgacgggta cttaactaaa atgacttgca gatggtcacc cagcacaatc    1200
69 caatcactag tgggaagcac tgtgcagctg aggtatcaca ggcgcagcct gtattgtcct    1260
71 gatagtccat ctattcatcc tacgtctgag cccaaaaact gcgtcttaca gagagacggc    1320
73 ttttatgaat gtgttttcca gccaatcttt ctattatctg ctatacaat gtggatcagg    1380
75 atcaaccatt ctttaggttc acttgactcg ccaccaacgt gtgtccttcc tgactccgta    1440
77 gtaaaaccac tacctccatc taacgtaaaa gcagagatta ctgtaaacac tggattattg    1500
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 79 aaagtatctt gggaaaagcc agtctttccg gagaataacc ttcaattcca gattcgatat   1560
 81 ggcttaagtg gaaaagaaat acaatggaag acacatgagg tattcgatgc aaagtcaaag   1620
 83 tctgccagcc tgctggtgtc agacctctgt gcagtctatg tggtccaggt tcgctgccgg   1680
 85 cggttggatg gactaggata ttggagtaat tggagcagtc cagcctatac gcttgtcatg   1740
 87 gatgtaaaag ttcctatgag agggcctgaa tttttggagaa aaatggatgg ggacgttact   1800
 89 aaaaaggaga gaaatgtcac cttgctttgg aagcccctga cgaaaaatga ctcactgtgt   1860
 91 agtgtgagga ggtacgtggt gaagcatcgt actgcccaca atgggacgtg gtcagaagat   1920
 93 gtgggaaatc ggaccaatct cactttcctg tggacagaac cagcgcacac tgttacagtt   1980
 95 ctggctgtca attccctcgg cgcttccctt gtgaatttta accttacctt ctcatggccc   2040
 97 atgagtaaag tgagtgctgt ggagtcactc agtgcttatc ccctgagcag cagctgtgtc   2100
 99 atcctttcct ggacactgtc acctgatgat tatagtctgt tatatctggt tattgaatgg   2160
101 aagatcctta atgaagatga tggaatgaag tggcttagaa ttccctcgaa tgttaaaaag   2220
103 ttttatatcc acgataattt tattcccatc gagaaatatc agtttagtct ttacccagta   2280
105 tttatggaag gagttggaaa accaaagata attaatggtt tcaccaaaga tgctatcgac   2340
107 aagcagcaga atgacgcagg gctgtatgtc attgtaccca taattatttc ctcttgtgtc   2400
109 ctactgctcg gaacactgtt aatttcacac cagagaatga aaaagttgtt ttgggacgat   2460
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117 <211> LENGTH: 842
118 <212> TYPE: PRT
119 <213> ORGANISM: Mus musculus
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124 <223> OTHER INFORMATION: X can be any amino acid
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W--> **133 Gln Ala Leu Ser Pro Cys Arg Ile Ser Thr Ser Leu Xaa Leu Val Pro**

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134             20                  25                  30
137 Asn Ser Ala Arg Gly Cys Phe Gly Asn Glu Gln Gly Gln Asn Cys Ser
138             35                  40                  45
141 Ala Leu Thr Asp Asn Thr Glu Gly Lys Thr Leu Ala Ser Val Val Lys
142     50                  55                  60
145 Ala Ser Val Phe Arg Gln Leu Gly Val Asn Trp Asp Ile Glu Cys Trp
146 65                  70                  75                  80
149 Met Lys Gly Asp Leu Thr Leu Phe Ile Cys His Met Glu Pro Leu Pro
150             85                  90                  95
153 Lys Asn Pro Phe Lys Asn Tyr Asp Ser Lys Val His Leu Leu Tyr Asp
154             100                 105                 110
157 Leu Pro Glu Val Ile Asp Asp Ser Pro Leu Pro Pro Leu Lys Asp Ser
158     115                 120                 125
161 Phe Gln Thr Val Gln Cys Asn Cys Ser Leu Arg Gly Cys Glu Cys His
162     130                 135                 140
165 Val Pro Val Pro Arg Ala Lys Leu Asn Tyr Ala Leu Leu Met Tyr Leu
166 145                 150                 155                 160
169 Glu Ile Thr Ser Ala Gly Val Ser Phe Gln Ser Pro Leu Met Ser Leu
170             165                 170                 175
```

```
173 Gln Pro Met Leu Val Val Lys Pro Asp Pro Pro Leu Gly Leu His Met
174         180             185             190
177 Glu Val Thr Asp Asp Gly Asn Leu Lys Ile Ser Trp Asp Ser Gln Thr
178         195             200             205
181 Met Ala Pro Phe Pro Leu Gln Tyr Gln Val Lys Tyr Leu Glu Asn Ser
182     210             215             220
185 Thr Ile Val Arg Glu Ala Ala Glu Ile Val Ser Ala Thr Ser Leu Leu
186 225             230             235             240
189 Val Asp Ser Val Leu Pro Gly Ser Ser Tyr Glu Val Gln Val Arg Ser
190             245             250             255
193 Lys Arg Leu Asp Gly Ser Gly Val Trp Ser Asp Trp Ser Ser Pro Gln
194             260             265             270
197 Val Phe Thr Thr Gln Asp Val Val Tyr Phe Pro Pro Lys Ile Leu Thr
198         275             280             285
201 Ser Val Gly Ser Asn Ala Ser Phe His Cys Ile Tyr Lys Asn Glu Asn
202     290             295             300
205 Gln Ile Ile Ser Ser Lys Gln Ile Val Trp Trp Arg Asn Leu Ala Glu
206 305             310             315             320
209 Lys Ile Pro Glu Ile Gln Tyr Ser Ile Val Ser Asp Arg Val Ser Lys
210             325             330             335
213 Val Thr Phe Ser Asn Leu Lys Ala Thr Arg Pro Arg Gly Lys Phe Thr
214             340             345             350
217 Tyr Asp Ala Val Tyr Cys Cys Asn Glu Gln Ala Cys His His Arg Tyr
218         355             360             365
221 Ala Glu Leu Tyr Val Ile Asp Val Asn Ile Asn Ile Ser Cys Glu Thr
222     370             375             380
225 Asp Gly Tyr Leu Thr Lys Met Thr Cys Arg Trp Ser Pro Ser Thr Ile
226 385             390             395             400
229 Gln Ser Leu Val Gly Ser Thr Val Gln Leu Arg Tyr His Arg Arg Ser
230             405             410             415
233 Leu Tyr Cys Pro Asp Ser Pro Ser Ile His Pro Thr Ser Glu Pro Lys
234             420             425             430
237 Asn Cys Val Leu Gln Arg Asp Gly Phe Tyr Glu Cys Val Phe Gln Pro
238         435             440             445
241 Ile Phe Leu Leu Ser Gly Tyr Thr Met Trp Ile Arg Ile Asn His Ser
242     450             455             460
245 Leu Gly Ser Leu Asp Ser Pro Pro Thr Cys Val Leu Pro Asp Ser Val
246 465             470             475             480
249 Val Lys Pro Leu Pro Pro Ser Asn Val Lys Ala Glu Ile Thr Val Asn
250             485             490             495
253 Thr Gly Leu Leu Lys Val Ser Trp Glu Lys Pro Val Phe Pro Glu Asn
254             500             505             510
257 Asn Leu Gln Phe Gln Ile Arg Tyr Gly Leu Ser Gly Lys Glu Ile Gln
258         515             520             525
261 Trp Lys Thr His Glu Val Phe Asp Ala Lys Ser Lys Ser Ala Ser Leu
262     530             535             540
265 Leu Val Ser Asp Leu Cys Ala Val Tyr Val Val Gln Val Arg Cys Arg
266 545             550             555             560
269 Arg Leu Asp Gly Leu Gly Tyr Trp Ser Asn Trp Ser Ser Pro Ala Tyr
```

```
270                   565                  570                  575
273 Thr Leu Val Met Asp Val Lys Val Pro Met Arg Gly Pro Glu Phe Trp
274               580                  585                  590
277 Arg Lys Met Asp Gly Asp Val Thr Lys Lys Glu Arg Asn Val Thr Leu
278           595                  600                  605
281 Leu Trp Lys Pro Leu Thr Lys Asn Asp Ser Leu Cys Ser Val Arg Arg
282       610                  615                  620
285 Tyr Val Val Lys His Arg Thr Ala His Asn Gly Thr Trp Ser Glu Asp
286 625                  630                  635                  640
289 Val Gly Asn Arg Thr Asn Leu Thr Phe Leu Trp Thr Glu Pro Ala His
290               645                  650                  655
293 Thr Val Thr Val Leu Ala Val Asn Ser Leu Gly Ala Ser Leu Val Asn
294               660                  665                  670
297 Phe Asn Leu Thr Phe Ser Trp Pro Met Ser Lys Val Ser Ala Val Glu
298           675                  680                  685
301 Ser Leu Ser Ala Tyr Pro Leu Ser Ser Ser Cys Val Ile Leu Ser Trp
302       690                  695                  700
305 Thr Leu Ser Pro Asp Asp Tyr Ser Leu Leu Tyr Leu Val Ile Glu Trp
306 705                  710                  715                  720
309 Lys Ile Leu Asn Glu Asp Asp Gly Met Lys Trp Leu Arg Ile Pro Ser
310               725                  730                  735
313 Asn Val Lys Lys Phe Tyr Ile His Asp Asn Phe Ile Pro Ile Glu Lys
314               740                  745                  750
317 Tyr Gln Phe Ser Leu Tyr Pro Val Phe Met Glu Gly Val Gly Lys Pro
318           755                  760                  765
321 Lys Ile Ile Asn Gly Phe Thr Lys Asp Ala Ile Asp Lys Gln Gln Asn
322       770                  775                  780
325 Asp Ala Gly Leu Tyr Val Ile Val Pro Ile Ile Ile Ser Ser Cys Val
326 785                  790                  795                  800
329 Leu Leu Leu Gly Thr Leu Leu Ile Ser His Gln Arg Met Lys Lys Leu
330               805                  810                  815
333 Phe Trp Asp Asp Val Pro Asn Pro Lys Asn Cys Ser Trp Ala Gln Gly
334           820                  825                  830
337 Leu Asn Phe Gln Lys Arg Thr Asp Thr Leu
338           835                  840
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342 <211> LENGTH: 2848
343 <212> TYPE: DNA
344 <213> ORGANISM: Mus musculus
346 <220> FEATURE:
347 <221> NAME/KEY: misc_feature
348 <222> LOCATION: (44)..(44)
349 <223> OTHER INFORMATION: N can be A, C, T or G
352 <220> FEATURE:
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358 <220> FEATURE:
359 <221> NAME/KEY: misc_feature
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RAW SEQUENCE LISTING   DATE: 02/11/2003
PATENT APPLICATION: US/08/599,974E   TIME: 10:56:27

Input Set : A:\EP.txt
Output Set: N:\CRF4\02112003\H599974E.raw

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361 <223> OTHER INFORMATION: N can be A, C, T or G
364 <220> FEATURE:
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367 <223> OTHER INFORMATION: N can be A, C, T or G
370 <220> FEATURE:
371 <221> NAME/KEY: misc_feature
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373 <223> OTHER INFORMATION: N can be A, C, T or G
376 <220> FEATURE:
377 <221> NAME/KEY: misc_feature
378 <222> LOCATION: (564)..(564)
379 <223> OTHER INFORMATION: N can be A, C, T or G
382 <220> FEATURE:
383 <221> NAME/KEY: misc_feature
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385 <223> OTHER INFORMATION: N can be A, C, T or G
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390 <222> LOCATION: (1335)..(1335)
391 <223> OTHER INFORMATION: N can be A, C, T or G
394 <220> FEATURE:
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396 <222> LOCATION: (2038)..(2038)
397 <223> OTHER INFORMATION: N can be A, C, T or G
400 <220> FEATURE:
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402 <222> LOCATION: (2179)..(2179)
403 <223> OTHER INFORMATION: N can be A, C, T or G
406 <220> FEATURE:
407 <221> NAME/KEY: misc_feature
408 <222> LOCATION: (2182)..(2182)
409 <223> OTHER INFORMATION: N can be A, C, T or G
412 <220> FEATURE:
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432 <222> LOCATION: (2610)..(2610)
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RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 02/11/2003
PATENT APPLICATION: **US/08/599,974E**   TIME: 10:56:28

Input Set : **A:\EP.txt**
Output Set: **N:\CRF4\02112003\H599974E.raw**

Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:2; Xaa Pos. 29
Seq#:3; N Pos. 44,67,234,483,527,564,1237,1335,2038,2179,2182,2183,2219
Seq#:3; N Pos. 2676,2610
Seq#:4; Xaa Pos. 79
Seq#:5; N Pos. 160,258
Seq#:6; Xaa Pos. 14,19,25,58,67,68,84,86
Seq#:35; N Pos. 5
Seq#:39; N Pos. 55,62,72,143
Seq#:40; N Pos. 83,101,181
Seq#:41; N Pos. 193
Seq#:57; Xaa Pos. 29
Seq#:58; Xaa Pos. 29
Seq#:59; Xaa Pos. 29
Seq#:60; Xaa Pos. 79
Seq#:61; Xaa Pos. 14,19,25,58,67,68,84,86
Seq#:62; Xaa Pos. 79
Seq#:63; Xaa Pos. 14,19,25,58,67,68,84,86

Invalid Line Length:
The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 8